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$RXR-\alpha$	<u>ILE.AELAVEPKTETYVEANMGL.NPSSPNDPVTNIC.QAADKQLFTL</u>	
PAR	LCQLGKYTTNSSADHRVQLDLGLWDKFSELATK.CII.KI	
PR	IN.LLM.SIEPDV.IYAGHD.N.TKPDTSSSLLTSL.NQLGERQLLSV	
GCR	VS.LLE.VIBPEV.LYAGYD.S.SVPDSTWRIMTTL.NMLGGRQVIAA	
ER	SALLD.A.EPPI.LYSEYD.P.TRPFSEASMMGLLTN.LADRELVHM	
NR1 ₀₁₁	IILLVSDDHEGRAA.QKRLETLLEERESKAEKVLQF.DP.GTKNVTAL	207
	Δ Δ Δ	
$RXR-\alpha$	V.EWAKRIPH.FSELPLDDQVILLRAGWNELLIASFSHR.SIA	
PAR	V.EFAKRLPG.FTGLSIADQITLLKAACLDILMLRICTR.YTP	
PR	V.KWSKSLPG.FRNLHIDDQITLIQYSWM.SLMV.FGLGWR.SYK	
GCR	V.KWAKAIPG.FRNLHLDDQMTLLQYSWM.FLMA.FALGWR.SYR	
ER	I.NWAKRVPG.FVDLTLHDQVHLLECAWLEILMIGLVWR.SME	
NR1 ₀₁₁	LME.ARELEARVIILSASEDDAATVYRAAAM.LNMTGSGYVWLVGER	252
$RXR-\alpha$	VKDG.IL.LATG.LH.VHR.N (SEQ ID NO:1)	
PAR	EQDT.MT.FSDG.LT.LNR (SEQ ID NO:2)	
PR	HVSGQMLYFAPD.LI.LN (SEQ ID NO:3)	
GCR	QSSANLLCFAPD.LI.IN (SEQ ID NO:4)	
ER	H.PGKLL.FAPN.LL.LDR.N (SEQ ID NO:5)	
NR1 ₀₁₁	EISGNALRYAPDGIIGLQLIN 273 (SEQ ID NO:6)	

FIG. 23